Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- 1. (withdrawn) A polynucleotide comprising an endogenous variant of the nucleotide sequence of SEQ ID NO: 1, or a degenerate variant of said endogenous variant.
- 2. (withdrawn) A polynucleotide according to claim 1 further comprising a space variant.
- 3. (withdrawn) A polynucleotide encoding a \(\beta\)-amyloid peptide-binding (BBP) protein comprising a PXDGS motif beginning at amino acid 237.
- 4. (currently amended) An isolated protein comprising the amino acid of SEQ ID NO: 2.
- 5. (currently amended) An isolated protein comprising the amino acid of SEQ ID NO: 2 from amino acid 68 to amino acid 269.
- 6. (currently amended) An isolated protein comprising the amino acid sequence encoded by the cDNA insert of clone BBP1-fl deposited under accession number ATCC 98617.
- 7. (currently amended) An isolated protein comprising the amino acid sequence from amino acid 185 to amino acid 217 of SEQ ID NO: 2.
- 8. (currently amended) A <u>non-naturally occurring</u> fusion protein comprising <u>a BBP1 an</u> <u>amino acid sequence with homology of 90% or greater to SEQ ID NO: 2</u> linked to a <u>heterologous protein or peptide</u> sequence.
- 9. (cancelled)
- 10. (withdrawn) A method for determining a polynucleotide encoding a \(\beta\)-amyloid peptidebinding protein (BBP) in a sample comprising the steps of (a) hybridizing to a sample a probe specific for said polynucleotide under conditions effective for said probe to hybridize specifically to said polynucleotide; and (b) determining the hybridization of said probe to polynucleotides in the sample, wherein said probe comprises a nucleic acid sequence having a region of 20 or more base pairs at least 90% identical to the polynucleotide sequence of SEQ ID NO: 1.
- 11. (withdrawn) A method for determining a polynucleotide encoding a \(\beta\)-amyloid peptidebinding protein (BBP) in a sample comprising the steps of (a) hybridizing to a sample a probe specific for said polynucleotide under conditions effective for said probe to hybridize

specifically to said polynucleotide; and (b) determining the hybridization of said probe to polynucleotides in the sample, wherein said probe comprises a nucleic acid sequence having a region of 20 or more base pairs at least 90% identical to the polynucleotide sequence of the cDNA insert of ATCC 98617 or ATCC 98399.

- 12. (withdrawn) An antibody that binds specifically to a polypeptide comprising the amino acid sequence of SEQ ID NO:1.
- 13. (withdrawn) An antibody that binds specifically to a polypeptide comprising the amino acid sequence of the \(\beta\)-amyloid peptide binding protein encoded by the cDNA insert of ATCC 98617.
- 14. (withdrawn) An antibody that binds to an extracellular region of a BBP.
- 15. (withdrawn) An antibody according to claim 14 wherein the extracellular region comprises the PXDGS motif.
- 16. (withdrawn) A method for detecting in a sample a polypeptide comprising a region at least 90% identical to the amino acid sequence of SEQ ID NO: 2, said method comprising (a) incubating with a sample a reagent that bind specifically to said polypeptide under conditions effective for specific binding; and (b) determining the binding of said reagent to said polypeptide in the same.
- 17. (withdrawn) A method for detecting in a sample a polypeptide comprising a region at least 90% identical in sequence to the amino acid sequence of the \(\beta\)-amyloid peptide binding protein encoded by the cDNA insert of ATCC 98617, said method comprising (a) incubating with a sample a reagent that bind specifically to said polypeptide under conditions effective for specific binding; and (b) determining the binding of said reagent to said polypeptide in the same.
- 18. (withdrawn) A method for diagnosing a disease characterized by aberrant expression of human β-amyloid peptide (BAP), comprising (a) incubating a sample indicative of the aberrant expression of human β-amyloid peptide with a reagent comprising a polypeptide comprising a region at least 90% identical to the amino acid sequence of SEQ ID NO:2 under conditions effective for specific binding of said reagent to said human β-amyloid peptide in the sample.
- 19. (withdrawn) A method of diagnosing the disease characterized by aberrant expression of human β-amyloid peptide, comprising (a) incubating a sample indicative of the aberrant

expression of human β-amyloid peptide with a reagent comprising a polypeptide comprising a region at least 90% identical to the amino acid sequence of the β-amyloid peptide binding protein encoded by the cDNA insert of ATCC 98617 under conditions effective for specific binding of said reagent to said human β-amyloid peptide; and (b) determining the binding of said reagent to said human β-amyloid peptide in the sample.

- 20. (withdrawn) A diagnostic process comprising analyzing for the presence of a polynucleotide of claim 1 in a sample derived from a host.
- 21. (withdrawn) A method for identifying compounds which regulate the activity of a β-amyloid peptide binding protein comprising (a) incubating a sample comprising human β-amyloid peptide in a test medium containing said test compound and a reagent comprising a polypeptide comprising a region at least 90% identical to the amino acid sequence of SEQ ID NO:2 under conditions effective for specific binding of said reagent to said human β-amyloid peptide; (b) comparing the binding of said reagent to said peptide in the same in the presence and absence of said test compound; and (c) relating the difference between the binding in step (b) to the test compound regulating the activity of the β-amyloid peptide binding protein.
- 22. (withdrawn) A method for identifying compounds which regulate the activity of a β-amyloid peptide binding protein comprising (a) incubating a sample comprising human β-amyloid peptide in a test medium containing said test compound and a amino acid sequence of the β-amyloid peptide binding protein encoded by the cDNA insert of ATCC 98617 under conditions effective for specific binding of said reagent to said human β-amyloid peptide; (b) comparing the binding of said reagent to said peptide in the sample in the presence and absence of said test compound; and (c) relating the difference between the binding in step (b) to the test compound regulating the activity of the β-amyloid peptide peptide binding protein.
- 23. (withdrawn) A method for the treatment of a patient having need to inhibit β-amyloid peptide accumulation in the brain comprising administering to the patient a therapeutically effective amount of BBP1.
- 24. (withdrawn) A method for the treatment of a patient having need of such treatment comprising administering to the patient a therapeutically effective amount of an antibody which binds to an extracellular portion of BBP1.
- 25. (withdrawn) A transgenic or chimeric nonhuman animal comprising the polynucleotide of SEQ ID NO: 1 or a degenerate variant of said polynucleotide.

- 26. (withdrawn) The animal of claim 25 wherein the transgene is under the control of a regulatable expression system.
- 27. (withdrawn) A knockout human animal wherein at least one allele of the BBP1 gene has been functionally disrupted.
- 28. (withdrawn) A knockout human animal wherein at least one allele of the BBP1 gene can be functionally disrupted by the induction of the Cre gene.
- 29. (withdrawn) A knockout according to claim 28 wherein the Cre gene is under the control of a tissue specific promoter.
- 30. (withdrawn) A knockout according to claim 28 wherein the Cre gene is under the control of a developmentally specific promoter.
- 31. (withdrawn) A knockout according to claim 28 wherein the Cre gene is under the control of an inducible promoter.
- 32. (withdrawn) A method for inhibiting expression of the BBP1 gene comprising providing to a cell double stranded ribonucleic acid complementary to a portion of the BBP1 gene wherein said ribonucleic acid comprises about 600 base pairs.
- 33. (withdrawn) A method of inhibiting expression of the BBP1 gene in a cell comprising providing said cell with an antisense nucleic acid.
- 34. (new) The fusion protein of claim 8 wherein the protein sequence comprises a human ß-Amyloid Peptide (BAP).
- 35. (new) The fusion protein of claim 34 wherein the BAP is BAP42.
- 36. (new) The fusion protein of claim 8, wherein the protein sequence linked to SEQ ID NO: 2 comprises a heterologous protein.
- 37. (new) A fusion protein comprising the amino acid sequence of SEQ ID NO: 2 from amino acids 68 to 269 with two regions of sufficient length and hydrophobicity to transverse a cellular membrane as deposited under the accession number ATCC 98399.
- 38. (new) A non-naturally occurring fusion protein comprising the amino acid sequence of SEQ ID NO: 2 linked to a peptide sequence.
- 39. (new) The fusion protein of claim 8 wherein the protein linked to SEQ ID NO: 2 comprises maltose binding protein (MBP), glutathione-S-transferase (GST), or thioredoxin (TRX).